

SEQUENCE LISTING



- <110> Tew, Kenneth D.
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Chen, Zhijian
- <120> Nucleic Acid Encoding Human ABCA
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- <130> FCCC.99-08US
- <140> 10/088,467
- <141> 2002-06-24
- <150> PCT/US00/40789
- <151> 2000-08-31
- <150> 60/154,839
- <151> 1999-09-20
- <160> 36
- <170> FastSEQ for Windows Version 3.0

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- <211> 8040
- <212> DNA
- <213> Homo sapiens

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			740					745					750				
Trp	Phe	Ile	Thr	Gly	Phe	Val	Gln	Leu	Ser	Ile	Ser	Val	Thr	Ala	Leu		
		755					760					765					
Thr	Ala	Ile	Leu	Lys	Tyr	Gly	Gln	Val	Leu	Met	His	Ser	His	Val	Val		
	770					775					780						
Ile	Ile	Trp	Leu	Phe	Leu	Ala	Val	Tyr	Ala	Val	Ala	Thr	Ile	Met	Phe		
785					790					795					800		
Cys	Phe	Leu	Val	Ser	Val	Leu	Tyr	Ser	Lys	Ala	Lys	Leu	Ala	Ser	Ala		
			805						810					815			
Cys	Gly	Gly	Ile	Ile	Tyr	Phe	Leu	Ser	Tyr	Val	Pro	Tyr	Met	Tyr	Val		
			820					825					830				
Ala	Ile	Arg	Glu	Glu	Val	Ala	His	Asp	Lys	Ile	Thr	Ala	Phe	Glu	Lys		
		835					840					845					
Cys	Ile	Ala	Ser	Leu	Met	Ser	Thr	Thr	Ala	Phe	Gly	Leu	Gly	Ser	Lys		
	850					855					860						
Tyr	Phe	Ala	Leu	Tyr	Glu	Val	Ala	Gly	Val	Gly	Ile	Gln	Trp	His	Thr		
865					870					875					880		
Phe	Ser	Gln	Ser	Pro	Val	Glu	Gly	Asp	Asp	Phe	Asn	Leu	Leu	Leu	Ala		
			885					890						895			
Val	Thr	Met	Leu	Met	Val	Asp	Ala	Val	Val	Tyr	Gly	Ile	Leu	Thr	Trp		
		900						905					910				
Tyr	Ile	Glu	Ala	Val	His	Pro	Gly	Met	Tyr	Gly	Leu	Pro	Arg	Pro	Trp		
	915						920					925					
Tyr	Phe	Pro	Leu	Gln	Lys	Ser	Tyr	Trp	Leu	Gly	Ser	Gly	Arg	Thr	Glu		
	930					935					940						
Ala	Trp	Glu	Trp	Ser	Trp	Pro	Trp	Ala	Arg	Thr	Pro	Arg	Leu	Ser	Val		
945					950					955					960		
Met	Glu	Glu	Asp	Gln	Ala	Cys	Ala	Met	Glu	Ser	Arg	Arg	Phe	Glu	Glu		
			965						970					975			
Thr	Arg	Gly	Met	Glu	Glu	Glu	Pro	Thr	His	Leu	Pro	Leu	Val	Val	Cys		
		980						985					990				
Val	Asp	Lys	Leu	Thr	Lys	Val	Tyr	Lys	Asp	Asp	Lys	Lys	Leu	Ala	Leu		
		995					1000					1005					
Asn	Lys	Leu	Ser	Leu	Asn	Leu	Tyr	Glu	Asn	Gln	Val	Val	Ser	Phe	Leu		
	1010					1015					1020						
Gly	His	Asn	Gly	Ala	Gly	Lys	Thr	Thr	Thr	Met	Ser	Ile	Leu	Thr	Gly		
1025					1030					1035					1040		
Leu	Phe	Pro	Pro	Thr	Ser	Gly	Ser	Ala	Thr	Ile	Tyr	Gly	His	Asp	Ile		

Ser Gly Val Gly Ala Thr Cys Val Leu Lys Ser Pro Ala Asn Gly Ser
 1540 1545 1550
 Leu Gly Pro Thr Leu Asn Leu Ser Ser Gly Glu Ser Arg Leu Leu Ala
 1555 1560 1565
 Ala Arg Phe Phe Asp Ser Met Cys Leu Glu Ser Phe Thr Gln Gly Leu
 1570 1575 1580
 Pro Leu Ser Asn Phe Val Pro Pro Pro Pro Ser Pro Ala Pro Ser Asp
 1585 1590 1595 1600
 Ser Pro Ala Ser Pro Asp Glu Asp Leu Gln Ala Trp Asn Val Ser Leu
 1605 1610 1615
 Pro Pro Thr Ala Gly Pro Glu Met Trp Thr Ser Ala Pro Ser Leu Pro
 1620 1625 1630
 Arg Leu Val Arg Glu Pro Val Arg Cys Thr Cys Ser Ala Gln Gly Thr
 1635 1640 1645
 Gly Phe Ser Cys Pro Ser Ser Val Gly Gly His Pro Pro Gln Met Arg
 1650 1655 1660
 Val Val Thr Gly Asp Ile Leu Thr Asp Ile Thr Gly His Asn Val Ser
 1665 1670 1675 1680
 Glu Tyr Leu Leu Phe Thr Ser Asp Arg Phe Arg Leu His Arg Tyr Gly
 1685 1690 1695
 Ala Ile Thr Phe Gly Asn Val Leu Lys Ser Ile Pro Ala Ser Phe Gly
 1700 1705 1710
 Thr Arg Ala Pro Pro Met Val Arg Lys Ile Ala Val Arg Arg Ala Ala
 1715 1720 1725
 Gln Val Phe Tyr Asn Asn Lys Gly Tyr His Ser Met Pro Thr Tyr Leu
 1730 1735 1740
 Asn Ser Leu Asn Asn Ala Ile Leu Arg Ala Asn Leu Pro Lys Ser Lys
 1745 1750 1755 1760
 Gly Asn Pro Ala Ala Tyr Gly Ile Thr Val Thr Asn His Pro Met Asn
 1765 1770 1775
 Lys Thr Ser Ala Ser Leu Ser Leu Asp Tyr Leu Leu Gln Gly Thr Asp
 1780 1785 1790
 Val Val Ile Ala Ile Phe Ile Ile Val Ala Met Ser Phe Val Pro Ala
 1795 1800 1805
 Ser Phe Val Val Phe Leu Val Ala Glu Lys Ser Thr Lys Ala Lys His
 1810 1815 1820
 Leu Gln Phe Val Ser Gly Cys Asn Pro Ile Ile Tyr Trp Leu Ala Asn
 1825 1830 1835 1840
 Tyr Val Trp Asp Met Leu Asn Tyr Leu Val Pro Ala Thr Cys Cys Val
 1845 1850 1855
 Ile Ile Leu Phe Val Phe Asp Leu Pro Ala Tyr Thr Ser Pro Thr Asn
 1860 1865 1870
 Phe Pro Ala Val Leu Ser Leu Phe Leu Leu Tyr Gly Trp Ser Ile Thr
 1875 1880 1885
 Pro Ile Met Tyr Pro Ala Ser Phe Trp Phe Glu Val Pro Ser Ser Ala
 1890 1895 1900
 Tyr Val Phe Leu Ile Val Ile Asn Leu Phe Ile Gly Ile Thr Ala Thr
 1905 1910 1915 1920
 Val Ala Thr Phe Leu Leu Gln Leu Phe Glu His Asp Lys Asp Leu Lys
 1925 1930 1935
 Val Val Asn Ser Tyr Leu Lys Ser Cys Phe Leu Ile Phe Pro Asn Tyr
 1940 1945 1950
 Asn Leu Gly His Gly Leu Met Glu Met Ala Tyr Asn Glu Tyr Ile Asn
 1955 1960 1965
 Glu Tyr Tyr Ala Lys Ile Gly Gln Phe Asp Lys Met Lys Ser Pro Phe
 1970 1975 1980
 Glu Trp Asp Ile Val Thr Arg Gly Leu Val Ala Met Ala Val Glu Gly
 1985 1990 1995 2000
 Val Val Gly Phe Leu Leu Thr Ile Met Cys Gln Tyr Asn Phe Leu Arg
 2005 2010 2015
 Arg Pro Gln Arg Met Pro Val Ser Thr Lys Pro Val Glu Asp Asp Val

2020					2025					2030					
Asp	Val	Ala	Ser	Glu	Arg	Gln	Arg	Val	Leu	Arg	Gly	Asp	Ala	Asp	Asn
2035					2040					2045					
Asp	Met	Val	Lys	Ile	Glu	Asn	Leu	Thr	Lys	Val	Tyr	Lys	Ser	Arg	Lys
2050					2055					2060					
Ile	Gly	Arg	Ile	Leu	Ala	Val	Asp	Arg	Leu	Cys	Leu	Gly	Val	Arg	Pro
2065					2070					2075					2080
Gly	Glu	Cys	Phe	Gly	Leu	Leu	Gly	Val	Asn	Gly	Ala	Gly	Lys	Thr	Ser
2085					2090					2095					
Thr	Phe	Lys	Met	Leu	Thr	Gly	Asp	Glu	Ser	Thr	Thr	Gly	Gly	Glu	Ala
2100					2105					2110					
Phe	Val	Asn	Gly	His	Ser	Val	Leu	Lys	Glu	Leu	Leu	Gln	Val	Gln	Gln
2115					2120					2125					
Ser	Leu	Gly	Tyr	Cys	Pro	Gln	Cys	Asp	Ala	Leu	Phe	Asp	Glu	Leu	Thr
2130					2135					2140					
Ala	Arg	Glu	His	Leu	Gln	Leu	Tyr	Thr	Arg	Leu	Arg	Gly	Ile	Ser	Trp
2145					2150					2155					2160
Lys	Asp	Glu	Ala	Arg	Val	Val	Lys	Trp	Ala	Leu	Glu	Lys	Leu	Glu	Leu
2165					2170					2175					
Thr	Lys	Tyr	Ala	Asp	Lys	Pro	Ala	Gly	Thr	Tyr	Ser	Gly	Gly	Asn	Lys
2180					2185					2190					
Arg	Lys	Leu	Ser	Thr	Ala	Ile	Ala	Leu	Ile	Gly	Tyr	Pro	Ala	Phe	Ile
2195					2200					2205					
Phe	Leu	Asp	Glu	Pro	Thr	Thr	Gly	Met	Asp	Pro	Lys	Ala	Arg	Arg	Phe
2210					2215					2220					
Leu	Trp	Asn	Leu	Ile	Leu	Asp	Leu	Ile	Lys	Thr	Gly	Arg	Ser	Val	Val
2225					2230					2235					2240
Leu	Thr	Ser	His	Ser	Met	Glu	Glu	Cys	Glu	Ala	Leu	Cys	Thr	Arg	Leu
2245					2250					2255					
Ala	Ile	Met	Val	Asn	Gly	Arg	Leu	Arg	Cys	Leu	Gly	Ser	Ile	Gln	His
2260					2265					2270					
Leu	Lys	Asn	Arg	Phe	Gly	Asp	Gly	Tyr	Met	Ile	Thr	Val	Arg	Thr	Lys
2275					2280					2285					
Ser	Ser	Gln	Ser	Val	Lys	Asp	Val	Val	Arg	Phe	Phe	Asn	Arg	Asn	Phe
2290					2295					2300					
Pro	Glu	Ala	Met	Leu	Lys	Glu	Arg	His	His	Thr	Lys	Val	Gln	Tyr	Gln
2305					2310					2315					2320
Leu	Lys	Ser	Glu	His	Ile	Ser	Leu	Ala	Gln	Val	Phe	Ser	Lys	Met	Glu
2325					2330					2335					
Gln	Val	Ser	Gly	Val	Leu	Gly	Ile	Glu	Asp	Tyr	Ser	Val	Ser	Gln	Thr
2340					2345					2350					
Thr	Leu	Asp	Asn	Val	Phe	Val	Asn	Phe	Ala	Lys	Lys	Gln	Ser	Asp	Asn
2355					2360					2365					
Leu	Glu	Gln	Gln	Glu	Thr	Glu	Pro	Pro	Ser	Ala	Leu	Gln	Ser	Pro	Leu
2370					2375					2380					
Gly	Cys	Leu	Leu	Ser	Leu	Leu	Arg	Pro	Arg	Ser	Ala	Pro	Thr	Glu	Leu
2385					2390					2395					2400
Arg	Ala	Leu	Val	Ala	Asp	Glu	Pro	Glu	Asp	Leu	Asp	Thr	Glu	Asp	Glu
2405					2410					2415					
Gly	Leu	Ile	Ser	Phe	Glu	Glu	Glu	Arg	Ala	Gln	Leu	Ser	Phe	Asn	Thr
2420					2425					2430					
Asp	Thr	Leu	Cys												
2435															

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 <213> Artificial Sequence

 <220>

<223> PCR Primer
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<210> 4
 <211> 27
 <212> DNA
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<220>
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<400> 4
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 <213> Artificial Sequence

<220>
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<400> 5
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<220>
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ataagcttgc tgaggcggcg gagcgtggc 29

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acagcgattg catgacaggc ag 22

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agggagctgg ctacaccgac g 21

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<210> 26
<211> 188
<212> PRT
<213> Homo Sapiens

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1 5 10 15
Phe Leu Gly His Asn Gly Ala Gly Lys Thr Thr Thr Met Ser Ile Leu
20 25 30
Thr Gly Leu Phe Pro Pro Thr Ser Gly Thr Ala Tyr Ile Leu Gly Lys
35 40 45
Asp Ile Arg Ser Glu Met Ser Thr Ile Arg Gln Asn Leu Gly Val Cys
50 55 60
Pro Gln His Asn Val Leu Phe Asp Met Leu Thr Val Glu Glu His Ile
65 70 75 80
Trp Phe Tyr Ala Arg Leu Lys Gly Leu Ser Glu Lys His Val Lys Ala
85 90 95
Glu Met Glu Gln Met Ala Leu Asp Val Gly Leu Pro Ser Ser Lys Leu
100 105 110
Lys Ser Lys Thr Ser Gln Leu Ser Gly Gly Met Gln Arg Lys Leu Ser
115 120 125
Val Ala Leu Ala Phe Val Gly Gly Ser Lys Val Val Ile Leu Asp Glu
130 135 140
Pro Thr Ala Gly Val Asp Pro Tyr Ser Arg Arg Gly Ile Trp Glu Leu
145 150 155 160
Leu Leu Lys Tyr Arg Gln Gly Arg Thr Ile Ile Leu Ser Thr His His
165 170 175
Met Asp Glu Ala Asp Val Leu Gly Asp Arg Ile Ala
180 185

<210> 27

<211> 183
 <212> PRT
 <213> Homo Sapiens

<400> 27
 Ala Val Asp Arg Ile Cys Val Gly Ile Pro Pro Gly Glu Cys Phe Gly
 1 5 10
 Leu Leu Gly Val Asn Gly Ala Gly Lys Ser Ser Thr Phe Lys Met Leu
 20 25 30
 Thr Gly Asp Thr Thr Val Thr Arg Gly Asp Ala Phe Leu Asn Arg Asn
 35 40 45
 Ser Ile Leu Ser Asn Ile His Glu Val His Gln Asn Met Gly Tyr Cys
 50 55 60
 Pro Gln Phe Asp Ala Ile Thr Glu Leu Leu Thr Gly Arg Glu His Val
 65 70 75 80
 Glu Phe Phe Ala Leu Leu Arg Gly Val Pro Glu Lys Glu Val Gly Lys
 85 90 95
 Val Gly Glu Trp Ala Ile Arg Lys Leu Gly Leu Val Lys Tyr Gly Glu
 100 105 110
 Lys Tyr Ala Gly Asn Tyr Ser Gly Gly Asn Lys Arg Lys Leu Ser Thr
 115 120 125
 Ala Met Ala Leu Ile Gly Gly Pro Pro Val Val Phe Leu Asp Glu Pro
 130 135 140
 Thr Thr Gly Met Asp Pro Lys Ala Arg Arg Phe Leu Trp Asn Cys Ala
 145 150 155 160
 Leu Ser Val Val Lys Glu Gly Arg Ser Val Val Leu Thr Ser His Ser
 165 170 175
 Met Glu Glu Cys Glu Ala Leu
 180

<210> 28
 <211> 187
 <212> PRT
 <213> Mus musculus

<400> 28
 Ala Leu Asn Lys Leu Ser Leu Asn Leu Tyr Glu Asn Gln Val Val Ser
 1 5 10 15
 Phe Leu Gly His Asn Gly Ala Gly Lys Thr Thr Thr Met Ser Ile Leu
 20 25 30
 Thr Gly Leu Phe Pro Pro Thr Ser Gly Ser Ala Thr Ile Tyr Gly His
 35 40 45
 Asp Ile Arg Thr Glu Met Asp Glu Ile Arg Lys Asn Leu Gly Met Cys
 50 55 60
 Pro Gln His Asn Val Leu Phe Asp Arg Leu Thr Val Glu Glu His Leu
 65 70 75 80
 Trp Phe Tyr Ser Arg Leu Lys Ser Met Ala Gln Glu Glu Ile Arg Lys
 85 90 95
 Glu Thr Asp Lys Met Ile Glu Asp Leu Glu Leu Ser Asn Lys Arg His
 100 105 110
 Ser Leu Val Gln Thr Leu Ser Gly Gly Met Lys Arg Lys Leu Ser Val
 115 120 125
 Ala Ile Ala Phe Val Gly Gly Ser Arg Ala Ile Ile Leu Asp Glu Pro
 130 135 140
 Thr Ala Gly Val Asp Pro Tyr Ala Arg Arg Ala Ile Trp Asp Leu Ile
 145 150 155 160
 Leu Lys Tyr Lys Pro Gly Arg Thr Ile Leu Leu Ser Thr His His Met
 165 170 175
 Asp Glu Ala Asp Leu Leu Gly Asp Arg Ile Ala
 180 185

<210> 29
 <211> 184
 <212> PRT
 <213> Mus musculus

<400> 29
 Ala Val Asp Arg Leu Cys Leu Gly Val Cys Val Pro Gly Glu Cys Phe
 1 5 10 15
 Gly Leu Leu Gly Val Asn Gly Ala Gly Lys Thr Ser Thr Phe Lys Met
 20 25 30
 Leu Thr Gly Asp Glu Ser Thr Thr Gly Gly Glu Ala Phe Val Asn Gly
 35 40 45
 His Ser Val Leu Lys Asp Leu Leu Gln Val Gln Gln Ser Leu Gly Tyr
 50 55 60
 Cys Pro Gln Phe Asp Ala Leu Phe Asp Glu Leu Thr Ala Arg Glu His
 65 70 75 80
 Leu Gln Leu Tyr Thr Arg Leu Arg Gly Ile Pro Trp Lys Asp Glu Ala
 85 90 95
 Gln Val Val Lys Trp Ala Leu Glu Lys Leu Glu Leu Thr Lys Tyr Ala
 100 105 110
 Asp Lys Pro Ala Gly Thr Tyr Ser Gly Gly Asn Lys Arg Lys Leu Ser
 115 120 125
 Thr Ala Ile Ala Leu Ile Gly Tyr Pro Ala Phe Ile Phe Leu Asp Glu
 130 135 140
 Pro Thr Thr Gly Met Asp Pro Lys Ala Arg Arg Phe Leu Trp Asn Leu
 145 150 155 160
 Ile Leu Asp Leu Ile Lys Thr Gly Arg Ser Val Val Leu Thr Ser His
 165 170 175
 Ser Met Glu Glu Cys Glu Ala Leu
 180

<210> 30
 <211> 187
 <212> PRT
 <213> Homo sapiens

<400> 30
 Ala Leu Asn Lys Leu Ser Leu Asn Leu Tyr Glu Asn Gln Val Val Ser
 1 5 10 15
 Phe Leu Gly His Asn Gly Ala Gly Lys Thr Thr Thr Met Ser Ile Leu
 20 25 30
 Thr Gly Leu Phe Pro Pro Thr Ser Gly Ser Ala Thr Ile Tyr Gly His
 35 40 45
 Asp Ile Arg Thr Glu Met Asp Glu Ile Arg Lys Asn Leu Gly Met Cys
 50 55 60
 Pro Gln His Asn Val Leu Phe Asp Arg Leu Thr Val Glu Glu His Leu
 65 70 75 80
 Trp Phe Tyr Ser Arg Leu Lys Ser Met Ala Gln Glu Glu Ile Arg Arg
 85 90 95
 Glu Met Asp Lys Met Ile Glu Asp Leu Glu Leu Ser Asn Lys Arg His
 100 105 110
 Ser Leu Val Gln Thr Leu Ser Gly Gly Met Lys Arg Lys Leu Ser Val
 115 120 125
 Ala Ile Ala Phe Val Gly Gly Ser Arg Ala Ile Ile Leu Asp Glu Pro
 130 135 140
 Thr Ala Gly Val Asp Pro Tyr Ala Arg Arg Ala Ile Trp Asp Leu Ile
 145 150 155 160
 Leu Lys Tyr Lys Pro Gly Arg Thr Ile Leu Leu Ser Thr His His Met

				165					170						175
Asp	Glu	Ala	Asp	Leu	Leu	Gly	Asp	Arg	Ile	Ala					
			180					185							

<210> 31
 <211> 183
 <212> PRT
 <213> Homo sapiens

<400> 31

Ala	Val	Asp	Arg	Leu	Cys	Leu	Gly	Val	Arg	Pro	Gly	Glu	Cys	Phe	Gly
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Leu	Leu	Gly	Val	Asn	Gly	Ala	Gly	Lys	Thr	Ser	Thr	Phe	Lys	Met	Leu
			20					25					30		
Thr	Gly	Asp	Glu	Ser	Thr	Thr	Gly	Gly	Glu	Ala	Phe	Val	Asn	Gly	His
			35				40					45			
Ser	Val	Leu	Lys	Glu	Leu	Leu	Gln	Val	Gln	Gln	Ser	Leu	Gly	Tyr	Cys
			50			55					60				
Pro	Gln	Cys	Asp	Ala	Leu	Phe	Asp	Glu	Leu	Thr	Ala	Arg	Glu	His	Leu
65					70					75					80
Gln	Leu	Tyr	Thr	Arg	Leu	Arg	Gly	Ile	Ser	Trp	Lys	Asp	Glu	Ala	Arg
				85				90						95	
Val	Val	Lys	Trp	Ala	Leu	Glu	Lys	Leu	Glu	Leu	Thr	Lys	Tyr	Ala	Asp
			100					105					110		
Lys	Pro	Ala	Gly	Thr	Tyr	Ser	Gly	Gly	Asn	Lys	Arg	Lys	Leu	Ser	Thr
			115				120					125			
Ala	Ile	Ala	Leu	Ile	Gly	Tyr	Pro	Ala	Phe	Ile	Phe	Leu	Asp	Glu	Pro
					135						140				
Thr	Thr	Gly	Met	Asp	Pro	Lys	Ala	Arg	Arg	Phe	Leu	Trp	Asn	Leu	Ile
145					150					155					160
Leu	Asp	Leu	Ile	Lys	Thr	Gly	Arg	Ser	Val	Val	Leu	Thr	Ser	His	Ser
				165					170					175	
Met	Glu	Glu	Cys	Glu	Ala	Leu									
			180												

<210> 32
 <211> 187
 <212> PRT
 <213> Homo sapiens

<400> 32

Ala	Val	Arg	Asp	Leu	Asn	Leu	Asn	Leu	Tyr	Glu	Gly	Gln	Ile	Thr	Val
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Leu	Leu	Gly	His	Asn	Gly	Ala	Gly	Lys	Thr	Thr	Thr	Leu	Ser	Met	Leu
			20					25					30		
Thr	Gly	Leu	Phe	Pro	Pro	Thr	Ser	Gly	Arg	Ala	Tyr	Ile	Ser	Gly	Tyr
			35				40					45			
Glu	Ile	Ser	Gln	Asp	Met	Val	Gln	Ile	Arg	Lys	Ser	Leu	Gly	Leu	Cys
			50			55					60				
Pro	Gln	His	Asp	Ile	Leu	Phe	Asp	Asn	Leu	Thr	Val	Ala	Glu	His	Leu
65					70					75					80
Tyr	Phe	Tyr	Ala	Gln	Leu	Lys	Gly	Leu	Ser	Arg	Gln	Lys	Cys	Pro	Glu
				85				90						95	
Glu	Val	Lys	Gln	Met	Leu	His	Ile	Ile	Gly	Leu	Glu	Asp	Lys	Trp	Asn
			100					105					110		
Ser	Arg	Ser	Arg	Phe	Leu	Ser	Gly	Gly	Met	Arg	Arg	Lys	Leu	Ser	Ile
			115				120					125			
Gly	Ile	Ala	Leu	Ile	Ala	Gly	Ser	Lys	Val	Leu	Ile	Leu	Asp	Glu	Pro
						135						140			

Thr	Ser	Gly	Met	Asp	Ala	Ile	Ser	Arg	Arg	Ala	Ile	Trp	Asp	Leu	Leu
145					150					155					160
Gln	Arg	Gln	Lys	Ser	Asp	Arg	Thr	Ile	Val	Leu	Thr	Thr	His	Phe	Met
				165					170					175	
Asp	Glu	Ala	Asp	Leu	Leu	Gly	Asp	Arg	Ile	Ala					
			180					185							

<210> 33
 <211> 183
 <212> PRT
 <213> Homo sapiens

<400> 33															
Ala	Val	Asp	Arg	Leu	Ser	Leu	Ala	Val	Gln	Lys	Gly	Glu	Cys	Phe	Gly
1				5					10					15	
Leu	Leu	Gly	Phe	Asn	Gly	Ala	Gly	Lys	Thr	Thr	Thr	Phe	Lys	Met	Leu
			20					25					30		
Thr	Gly	Glu	Glu	Ser	Leu	Thr	Ser	Gly	Asp	Ala	Phe	Val	Gly	Gly	His
		35					40					45			
Arg	Ile	Ser	Ser	Asp	Val	Gly	Lys	Val	Arg	Gln	Arg	Ile	Gly	Tyr	Cys
	50				55						60				
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Val	Met	Tyr	Ala	Arg	Leu	Arg	Gly	Ile	Pro	Glu	Arg	His	Ile	Gly	Ala
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Cys	Val	Glu	Asn	Thr	Leu	Arg	Gly	Leu	Leu	Leu	Glu	Pro	His	Ala	Asn
			100					105					110		
Lys	Leu	Val	Arg	Thr	Tyr	Ser	Gly	Gly	Asn	Lys	Arg	Lys	Leu	Ser	Thr
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Gly	Ile	Ala	Leu	Ile	Gly	Glu	Pro	Ala	Val	Ile	Phe	Leu	Asp	Glu	Pro
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Ser	Thr	Gly	Met	Asp	Pro	Val	Ala	Arg	Arg	Leu	Leu	Trp	Asp	Thr	Val
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Ala	Arg	Ala	Arg	Glu	Ser	Gly	Lys	Ala	Ile	Ile	Ile	Thr	Ser	His	Ser
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Thr	Gly	Leu	Leu	Pro	Pro	Thr	Ser	Gly	Thr	Val	Leu	Val	Gly	Gly	Arg
		35					40					45			
Asp	Ile	Glu	Thr	Ser	Leu	Asp	Ala	Val	Arg	Gln	Ser	Leu	Gly	Met	Cys
	50				55						60				
Pro	Gln	His	Asn	Ile	Leu	Phe	His	His	Leu	Thr	Val	Ala	Glu	His	Met
65					70					75					80
Leu	Phe	Tyr	Ala	Gln	Leu	Lys	Gly	Lys	Ser	Gln	Glu	Glu	Ala	Gln	Leu
				85					90					95	
Glu	Met	Glu	Ala	Met	Leu	Glu	Asp	Thr	Gly	Leu	His	His	Lys	Arg	Asn
			100					105					110		
Glu	Glu	Ala	Gln	Asp	Leu	Ser	Gly	Gly	Met	Gln	Arg	Lys	Leu	Ser	Val

